SEQUENCE LISTING

SEQ 1: Arabidopsis thaliana GAD1

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SEQ2: Arabidopsis thaliana GAD1

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SEQ 3: Arabidopsis thaliana GAD2

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SEQ 4: Arabidopsis thaliana GAD2

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SEQ 5: Arabidopsis thaliana GAD3

ATGGTTTTATCTAAGACAGCTTCCAAATCCGATGATTCAATCCATTCAACTTTTGCTTCCCGTTATGTC ${\tt CGCAACTCTATCTCACGGTAAGAAGTTGAAACACAATTTTATTTTGTTTAATGTTTTCATTGGTAACTA}$ ${\tt TCTAACGAAATTAACCACTTGCACTGAAAGATCCGAGCATAATGTGTGTTACTATATAAGAGGTATTTT}$ ${\tt CTTTTTTAATCTTAAGCTAAATATATCAATTTTTCATCAGATTCGAAATACCTAAGAACTCGATCCCTA}$ AGGAAGCAGCATACCAAATCATCAACGACGAGCTCAAGTTTGACGGTAACCCGAGGCTAAACCTGGCCT CCTTTGTGACCACTTGGATGGAGCCAGAATGTGACAAGCTCATGATGGAATCCATCAACAAGAACAACG TTGAGATGGACCAATACCCTGTTACCACCGACCTTCAGAATCGATGCGTTAACATGATTGCGCGTCTCT TCAACGCGCCTTTAGGTGACGGTGAAGCCGCCATTGGTGTTGGCACGGTGGGGTCATCGGAGGCAGTGA TGTTGGCCGGACTGGCCTTTAAGAGACAGTGGCAGAACAAGCGTAAGGCCCTAGGGCTGCCTTATGATA GACCTAATATTGTAACCGGAGCCAATATTCAGGTAAACCAAAACAAAATTGATTAAATTTTAAACCGG $\tt TTTAGGTCTATGTTTACATTGACTCAATTTCCGGTTCAATACAGGTTTGCTTGGAGAAATTTGCAAGGT$ ATTTTGAAGTGGAGCTTAAGGAAGTGAAGCTGAGAGAAGGATATTACGTGATGGACCCTGACAAAGCGG TTGAAATGGTAGACGAAAACACTATATGCGTCGTGGCCATCCTCGGTTCGACACTAACCGGAGAATTCG ${\tt AAGACGTTAAGCTCCTCAACGACCTTTTAGTCGAGAAAAACAAGAAAACCGGGTAATTGAATCAAAACC}$ ${\tt AACTAACAAATTAATTTTATATACTTTTGCCTAGAAATATTACAATTTCTAACGTGAGATATATTTGCT}$ ${\tt TAGAAATATTTTTTTTTTTTTTTAATGAATAAAACTTATTAACCAAAACAAAACCATATTTTACATT}$ ${ t ATATGCTTCCTTGTATCGAATGGTGTTTTAAATACTGATTAAAAAATGTTTTGCTTAAAAAATATAACAA}$ ${\tt TTTATAATGTGAGATATTCAAGCATTCTAATATCAAACCGATAAACAACAACAAACTGATTATTAATTT}$ ${ t ATTTAACCGGTTTGGTTCCGGTTTAATATTTTGTAGATGGGATACGCCGATTCACGTGGACGCAGCGA}$ $\tt GTGGTGGGTTTATTGCTCCCTTCTTGTATCCGGACTTGGAGTGGGATTTCCGGTTACCGTTGGTTAAGA$ ${\tt GCATAAATGTGAGTGACAAATACGGTTTGGTTTACGCCGGTATCGGTTGGGTCGTATGGAGAACCA}$ $A \texttt{A} \texttt{A} \texttt{C} \texttt{C} \texttt{G} \texttt{A} \texttt{T} \texttt{G} \texttt{A} \texttt{C} \texttt{C} \texttt{A} \texttt{T} \texttt{A} \texttt{T} \texttt{C} \texttt{T} \texttt{T} \texttt{C} \texttt{C} \texttt{A} \texttt{T} \texttt{T} \texttt{A} \texttt{C} \texttt{C} \texttt{C} \texttt{A} \texttt{C} \texttt{T} \texttt{T} \texttt{A} \texttt{C} \texttt{C} \texttt{C} \texttt{C} \texttt{A} \texttt{T} \texttt{T} \texttt{A} \texttt{C} \texttt{C} \texttt{C} \texttt{C} \texttt{A} \texttt{C} \texttt{T} \texttt{T} \texttt{A} \texttt{C} \texttt{C} \texttt{C} \texttt{C} \texttt{A} \texttt{$ TCAACTTCTCTAAAGGTACATTACCATATCTTATGTAAAGTTTAGATATATTTATAGATTAATGTTTTG ${\tt TTAATTCTTGTATATTACCAGGGTCAAGTCAAGTGATTGCTCAGTACTACCAGTTGATTCGTCTTGGAT}$ ${\tt TCGAGGTAAATAACTCAATAAAGAAACTAAAACGTTACTAAATCCAATCGTATACGTACTAGTATA}$ ${\tt ATATACAAGTTGTTACTATACTTTATGACTACAAAAGTTCAAAAACCAAGAATGTACTAAATACATTCCA}$

TAGGGATATCGCAACGTGATGGATAATTGCCGCGAGAACATGATGGTACTAAGACAAGGATTAGAGAAA ACGGGACGTTTTAACATCGTCTCCAAAGAAAACGGTGTTCCGTTAGTGGCGTTTTCTCTCAAAGATAGT AGCCGCCACAACGAGTTCGAGGTGGCCGAAATGCTTCGTCGCTTCGGCTGGATCGTTCCGGCCTACACG ATGCCTGCGGATGCGCAACATGTCACGGTCCTTCGAGTTGTTATCCGAGAAGATTTCTCTCGAACCTTA GCTGAGAGATTGGTAGCCGATTCGAGAGGTTCTACACGAGCTCGATACGCTTCCGCGAGGGTTCAC GCCAAGATGGCTAGTGGAAAAGTTAACGGTGTTAAGAAGACGCCAGAGGAGACGCAAAGAAGAAGTCACG GCCTACTGGAAGAAGTTTGTGGACACTAAGACTGACAAGAACGGCGTTCCGTTAGTAGCAAGTATTACC AATCAATGA

SEQ 6: Arabidopsis thaliana GAD3

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SEQ 7: Arabidopsis thaliana GAD4

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SEQ 8: Arabidopsis thaliana GAD4

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 ${\tt NCRENMMVLRQGLEKTGRFKIVSKENGVPLVAFSLKDSSRHNEFEVAHTLRRFGWIVPAYTMPADAQHV} \\ {\tt TVLRVVIREDFSRTLAERLVADFEKVLHELDTLPARVHAKMANGKVNGVKKTPEETQREVTAYWKKLLE} \\ {\tt TKKTNKNTIC}$

SEQ 9: Arabidopsis thaliana GAD5

ATGGTACTCGCAACCAACTCTGACTCCGACGAGCATTTGCATTCCACTTTTGCTTCTAGATATGTCCGT ${\tt GCTGTTGTTCCCAGGGTTCCAGAGAGTTTTGCCTCATTTTAGTTTTTTTAATCTTGTATGCTACATTGTT}$ ${ t ATATATTTAATTTATGTATCTGTTTGCATATATTGAAACAGGTTCAAGATGCCTGACCATTGCATG$ ${\tt CCCAAAGATGCTGCTTATCAAGTGATCAATGATGAGTTGATGCTTGATGGTAATCCCAGGCTTAACCTA}$ GCCTCCTTTGTCACCACTTGGATGGAACCTGAGTGTGACAAACTCATCATGGATTCTGTCAATAAGAAC TATGTTGATATGGATGAATATCCTGTCACCACTGAGCTCCAGGTTCCTCCTTCTTTCCTCATTCTCT ${\tt CTCTCATCTACTTTCCACTGTTTTGTCATAGACTCATACATCTTTTATCTGGCTTATTTTTCAGAACCG}$ $\tt GTGTGTAAATATGATAGCAAACTTGTTCCATGCTCCCGTTGGAGAAGACGAGGCTGCTATTGGGTGTGG$ AACTGTTGGTTCATCTGAGGCTATAATGCTTGCTGGTTTGGCTTTCAAAAGGAAATGGCAACATAGGAG AAAAGCTCAGGGTCTACCTATTGATAAGCCTAACATTGTCACTGGAGCCAATGTTCAGGTCTAAAATAT ACATTTTCAAACTTTGTTTCATCCGCTCAGGTGTGCTGGGAGAGTTTGCAAGGTACTTTGAGGTAGAG $\tt CTCAAAGAGTGAAACTAAGTGAAGACTACTATGTTATGGATCCAGCTAAAGCTGTAGAGATGGTGGAT$ GAGAATACCATCTGTGTTGCAGCAATTCTAGGATCCACACTTACTGGAGAGTTTGAGGACGTTAAGCAA TTGAACGATCTCTTAGCTGAGAAAAACGCAGAGACAGGATGGGAAACTCCTATTCATGTTGATGCAGCC AGTGGAGGATTCATTGCTCCTTTCCTCTACCCTGATCTTGAATGGGACTTTAGGCTTCCATGGGTGAAG AAAGATGATTTGCCAGAGGAACTTGTCTTCCACATCAACTACTTGGGAGCTGATCAACCCACTTTCACT $\tt CTCAACTTCTCAAAAGGTTTGTAAAATAAAAACTGGCTTTATCCAATCAAATCCATCATCACATTTCCT$ ${\tt TTAAGAAACTCAATGTTTTCTTTTGCAGGGTCGAGCCAAATCATTGCTCAGTACTATCAGTTTATCCGA}$ $\tt CTAGGCTTTGAGGTACTTGTTCCCTTATCTGCATTACAGTTTCATTTTTCATCTTGCTTAATCTAATG$ TAAGAGAAGGAATAGAGATGACAGGGAAGTTCAACATTGTGTCCAAAGATATTGGCGTGCCACTAGTGG ${\tt CATTCTCTCAAAGACAGTAGCAAGCACACGGTGTTTGAGATCGCAGAGTCTTTGAGAAAATTCGGGT}$ GGATCATACCGGCTTACACTATGCCTGCAGATGCACAGCACATTGCTGTGCTCAGAGTTGTGATAAGAG GGCTTCCTAGCAGGATTGCACATCTTGCTGCGGCTGCAGCGGTTAGTGGTGATGATGAAGAAGTTAAAG ATATTGTCTGCTAA

SEQ 10: Arabidopsis thaliana GAD5

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SEQ 11: Tobacco NtGAD1

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SEQ 12: Tobacco NtGAD1

MVLSKTASESDVSIHSTFASRYVRTSLPRFKMPENSIPKEAAYQIINDELMLDGNPRLNLASFVTTWME PECNKLMMDSINKNYVDMDEYPVTTELQNRCVNMIAHLFNAPLGDGETAVGVGTVGSSEAIMLAGLAFK RKWQNKMKAQGKPCDKPNIVTGANVQVCWEKFARYFEVELKEVKLSDGYYVMDPEKAVEMVDENTICVA AILGSTLNGEFEDVKRLNDLLIEKNKETGWDTPIHVDAASGGFIAPFLYPELEWDFRLPLVKSINVSGH KYGLVYAGIGWAIWRNKEDLPDELIFHINYLGADQPTFTLNFSKGSSQVIAQYYQLIRLGFEGYKNVME NCQENARVLREGLEKSGRFNIISKEIGVPLVAFSLKDNSQHNEFEISETLRRFGWIIPAYTMPPNAQHV TVLRVVIREDFSRTLAERLVIDIEKVLHELDTLPARVNAKLAVAEANGSGVHKKTDREVQLEITTAWKK FVADKKKKTNGVC

SEQ 13: Tobacco NtGAD2

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1501 actgcatggt tgaaatttgt tgctgataag aagaagaaga ctaatggagt ttgttaattt
1561 aatttaacaa aaaaaaagtt tataatatgg tgatttatgt aactactagc agtcgtactg
1621 cttgtttttt atatttgagt tgatgtgttt tttgagcact tgaggagcta gctagttatt
1681 gctagtgaaa aattggatga tatattttgg actactttgt aagtttgtat tattaatcca
1741 aattaaacga tatttatcat aaaaaaaaa a
```

SEQ 14: Tobacco NtGAD2

MVLSKTASESDVSVHSTFASRYVRTSLPRFKMPENSIPKEAAYQIINDELMLDGNPRLNLASFVTTWME PECNTLMMDSINKNYVDMDEYPVTTELQNRCVNMIAHLFNAPLGDGETAVGVGTVGSSEAIMLAGLAFK RKWQNKMKAQGKPFDKPNIVTGANVQVCWEKFARYFEVELKEVKLSDGYYVMDPEKAVEMVDENTICVA AILGSTLNGEFEDVKRLNDLLIEKNKETGWDTPIHVDAASGGFIAPFLYPELEWDFRLPLEKSINVSGH KYGLVYAGIGWAIWRNKEDLPDELIFHINYLGADQPTFTLNFSKGSSQVIAQYYQLIRLGFEGYKNVME NCQENARVLREGIEKSGRFNIISKEIGVPLVAFSLKDNSQHNEFEISETLRRFGWIVLAYTMPPNAQHV TVLRVVIREDFSRTLAERLVIDIEKVFHGVDTLPARVNAKLAVAEANGSGVHKKTDREVQLEITTAWLK FVADKKKKTNGVC

SEQ 15: Petunia GAD

```
aaagagtaca aactaatatc cacttaaatt gtatttctcc attttctctc tttatttagt
1
61
     ctgtcataac aatggttcta tcaaagacag tgtcgcagag cgatgtgtcc attcactcca
     cgtttgcttc tcgatatgtt cgaacttctc ttcccaggtt taaaatgcca gataattcga
121
181
     taccaaaaga agcagcatat cagatcataa atgatgaact gatgttagat ggaaacccaa
     ggctgaactt ggcttctttt gttacaacat ggatggaacc agagtgtgat aagttgatga
241
301
     tggactctat taacaagaac tatgttgata tggatgaata tcctgttacc actgagcttc
361
     agaatcgatg tgtaaacatg atagctcatt tgtttaatgc accacttgaa gatggagaaa
421
     ctgcagttgg agttggaact gttggatcct ctgaagccat tatgcttgct ggattagctt
481
     tcaagagaaa atggcagaac aaaatgaaag cccaaggcaa accctgtgac aagcccaaca
    ttgttactgg tgcaaatgtc caggtgtgct gggagaaatt tgcaaggtat tttgaagtgg
541
    agctaaagga agtaaagctt agtgaaggat actatgtgat ggaccctgag aaagctgtgg
601
    agatggtgga tgaaaacacc atttgtgtag ctgctatctt aggttccacc ctcaatggag
661
    aatttgaaga cgttaagcgc ttgaatgatc tcttggtcga gaagaacaaa gaaaccgggt
721
781
    gggacactee aatteatgtg gatgeageaa gtggtggatt tattgeaceg tteatttace
841
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901 aatatggtct tgtctatgct ggtattggtt gggtcgtttg gaggaacaag atgatttgc
    ctgatgaact tatcttccac attaattatc ttggtgctga tcaacctact ttcactctca
1021 acttttctaa aggttctagc caagtaattg ctcaatatta ccaacttatt cgcttgggtt
1081 atgagggtta caagaatgtg atggagaatt gtcaagaaaa tgcatcggta ctaagagaag
1141 ggctagaaaa gacaggaaga ttcaacataa tctccaaaga aattggagta cctttagtag
1201 cattetetet taaagacaac aggeaacaca acgagttega gatttetgaa aetttaagga
1261 gatttggttg gattgttcct gcatatacta tgccaccaaa cgcacaacac attacagttc
1321 tcagagttgt gatcagagaa gatttctccc gtacgcttgc agaacgactg gtaagagaca
1381 tcgaaaaagt ccttcatgaa cttgacacac tccctgcacg tgtcaatgct aagctcgctg
1441 tggccgagga gcaggcggct gcgaatggca gcgaggtgca taagaaaaca gatagcgaag
1501 tgcagttgga gatgataact gcatggaaga agtttgttga agaaaagaag aagaagacta
1561 atcgagtttg ttaattaatt atattagtgt ttataatatg atgaatatgg ctattatcat
1621 tggtgactgc ttgttagtat attagctgtg attatcacca atatgagttt ggttttcttg
1681 atttggttct tttcagtact tgaaaagttg ttattgatat tgtaaaattg tacttttaa
1741 ctatttggat tattaatgcc aattttctag tgtacttaat aaaaa
```

SEQ 16: Petunia GAD

MVLSKTVSQSDVSIHSTFASRYVRTSLPRFKMPDNSIPKEAAYQIINDELMLDGNPRLNLASFVTTWME PECDKLMMDSINKNYVDMDEYPVTTELQNRCVNMIAHLFNAPLEDGETAVGVGTVGSSEAIMLAGLAFK RKWQNKMKAQGKPCDKPNIVTGANVQVCWEKFARYFEVELKEVKLSEGYYVMDPEKAVEMVDENTICVA AILGSTLNGEFEDVKRLNDLLVEKNKETGWDTPIHVDAASGGFIAPFIYPELEWDFRLPLVKSINVSGH KYGLVYAGIGWVVWRNKDDLPDELIFHINYLGADQPTFTLNFSKGSSQVIAQYYQLIRLGYEGYKNVME NCQENASVLREGLEKTGRFNIISKEIGVPLVAFSLKDNRQHNEFEISETLRRFGWIVPAYTMPPNAQHI TVLRVVIREDFSRTLAERLVRDIEKVLHELDTLPARVNAKLAVAEEQAAANGSEVHKKTDSEVQLEMIT AWKKFVEEKKKKTNRVC

SEQ 17: Tomato GAD

```
aaaaaatggt gttaacaacg acgtcgataa gagattcaga agagagcttg cactgtacat
61
     ttgcatcaag atatgtacag gaacctttac ctaagttcaa aatgcctaaa aaatccatgc
121
     cgaaagaagc agcttatcag attgtaaacg acgagcttat gttggatggt aaccccaggt
181
     tgaatttagc ttcctttgtt agcacatgga tggagcccga gtgcgataag ctcatcatgt
241
     catccattaa taaaaactat gtcgacatgg atgagtatcc tgtcaccact gaacttcaaa
301
     atagatgtgt taacatgtta gcacatcttt tccatgcccc ggttggtgat gatgagactg
361
     cagttggagt tggtacagtg ggttcatcag aggcaataat gcttgctggc cttgcttca
     aacgcaaatg gcaatcgaaa agaaaagcag aaggcaaacc tttcgataag cctaatatag
421
     tcactggagc taatgtgcag gtctgctggg aaaaatttgc aaggtatttt gaggttgagt
481
     tgaaggaggt gaaactaaaa gaaggatact atgtaatgga ccctgccaaa gcagtagaga
541
     tagtggatga gaatacaata tgtgttgctg caatccttgg ttctactctg actggggagt
601
     ttgaggatgt gaagctccta aacgagctcc ttacaaaaaa gaacaaggaa accggatggg
661
721
     agacaccgat tcatgtcgat gctgcgagtg gaggatttat tgctcctttc ctctggccag
781
     atcttgaatg ggatttccgt ttgcctcttg tgaaaagtat aaatgtcagc ggtcacaagt
841
     atggccttgt atatgctggt gtcggttggg tgatatggcg gagcaaggaa gacttgcccg
901
     atgaactcgt ctttcatata aactaccttg ggtctgatca gcctactttt actctcaact
961
     tctctaaagg ttcctatcaa ataattgcac agtattatca gttaataaga cttggctttg
1021 agggttataa gaacgtcatg aagaattgct tatcaaacgc aaaagtacta acagagggaa
1081 tcacaaaaat ggggcggttc gatattgtct ctaaggatgt gggtgttcct gttgtagcat
1141 tttctctcag ggacagcagc aaatatacgg tatttgaagt atctgagcat ctcagaagat
1201 ttggatggat cgtccctgca tacacaatgc caccggatgc tgaacacatt gctgtactgc
1261 gggttgtcat tagagaggat ttcagccaca gcctagctga gagacttgtt tctgacattg
1321 agaaaattct gtcagagttg gacacacagc ctcctcgttt gcccaccaaa gctgtccgtg
1381 tcactgctga ggaagtgcgt gatgacaagg gtgatgggct tcatcatttt cacatggata
1441 ctgtagagac tcagaaagac attatcaaac attggaggaa aatcgcaggg aagaagacca
1501 gcggagtctg ctaggtctgg ccacacttgt tatctgggct ccgcttccat cgccatcctg
1561 tagtatgtat tacgtgtgtt gtttccatct tatgtagtag ttggtactgt aatctgtgta
1621 aatgetttea tgatettgge tetgtatatg etaaataage actgeattte aagtteetgg
1681 aagtatttat gtatgaatca atccgggcat aattggtaga atgccctctc tgcgtcatct
1741 ttgaatttca cgtgcaataa tatttgaaat ctacacctat tat
```

SEQ 18: Tomato GAD

MVLTTTSIRDSEESLHCTFASRYVQEPLPKFKMPKKSMPKEAAYQIVNDELMLDGNPRLNLASFVSTWM EPECDKLIMSSINKNYVDMDEYPVTTELQNRCVNMLAHLFHAPVGDDETAVGVGTVGSSEAIMLAGLAF KRKWQSKRKAEGKPFDKPNIVTGANVQVCWEKFARYFEVELKEVKLKEGYYVMDPAKAVEIVDENTICV AAILGSTLTGEFEDVKLLNELLTKKNKETGWETPIHVDAASGGFIAPFLWPDLEWDFRLPLVKSINVSG HKYGLVYAGVGWVIWRSKEDLPDELVFHINYLGSDQPTFTLNFSKGSYQIIAQYYQLIRLGFEGYKNVM KNCLSNAKVLTEGITKMGRFDIVSKDVGVPVVAFSLRDSSKYTVFEVSEHLRRFGWIVPAYTMPPDAEH IAVLRVVIREDFSHSLAERLVSDIEKILSELDTQPPRLPTKAVRVTAEEVRDDKGDGLHHFHMDTVETQ KDIIKHWRKIAGKKTSGVC

1) $Arabidopsis\ thaliana\ ecotype\ Columbia\ glutamate\ decarboxylase\ 1\ (GAD1)\ cDNA$

Note: This is nucleic acid SEQ #1 and amino acid SEQ #2

A) LOCUS ATU10034 ACCESSION U10034 VERSION U10034.1 GI:497978 REFERENCE AUTHORS Arazi, T., Baum, G., Snedden, W.A., Shelp, B.J. and Fromm, H. TITLE Molecular and biochemical analysis of calmodulin interactions with the calmodulin-binding domain of plant glutamate decarboxylase JOURNAL Plant Physiol. 108 (2), 551-561 (1995)

1. From Arabidopsis genome sequencing project chromosome 5 (ACC# AB005238)

LOCUS BAB10520

DEFINITION glutamate decarboxylase 1 (GAD 1) (Arabidopsis thaliana)

ACCESSION BAB10520

PID g10177078

VERSION BAB10520.1 GI:10177078

REFERENCE 1 (sites)

AUTHORS Sato, S., Kotani, H., Nakamura, Y., Kaneko, T., Asamizu, E., Fukami, M., Miyajima, N. and Tabata, S.

TITLE Structural analysis of Arabidopsis thaliana chromosome 5. I. Sequence features of the 1.6 Mb regions covered by twenty physically assigned P1 clones

JOURNAL DNA Res. 4 (3), 215-230 (1997)

2) Arabidopsis thaliana ecotype Columbia glutamate decarboxylase 2 (GAD2) cDNA

Note: This is nucleic acid SEQ #3 and amino acid SEO #4

A) LOCUS ATU46665

ACCESSION U46665

VERSION U46665.1 GI:1184959

REFERENCE

AUTHORS Turano, F.J. and Fang, T.K.

TITLE Characterization of two glutamate decarboxylase cDNA clones from Arabidopsis

JOURNAL Plant Physiol. 117 (4), 1411-1421 (1998)

B) LOCUS ATU49937

ACCESSION U49937

VERSION U49937.1 GI:1236618

REFERENCE

AUTHORS Zik, M., Arazi, T., Snedden, W.A. and Fromm, H.

TITLE Two isoforms of glutamate decarboxylase in Arabidopsis a regulated by calcium/calmodulin and differ in organ distribution JOURNAL Plant Mol. Biol. 37 (6), 967-975 (1998)

C) From Arabidopsis genome sequencing project

ACCESSION #AC009513

Part of chromosome #1

note="Identical to gblU46665 glutamate decarboxylase 2 (GAD 2)

Arabidopsis thaliana. and ESTs gblW43856, gblN37724,

gblZ34642 and gblR90491 come from this gene."

/protein_id="AAF06056.1"

/db_xref="GI:6227020"

3) *Arabidopsis thaliana* ecotype Columbia putative glutamate decarboxylase (putative GAD3) DNA From Arabidopsis genome sequencing project

Note: This is nucleic acid SEQ #5 and amino acid SEQ #6

ACCESSION #AC006532

Part of chromosome #2 /product="putative glutamate decarboxylase" /protein_id="AAD20093.1" /db xref="GI:4406783"

4) *Arabidopsis thaliana* ecotype Columbia putative glutamate decarboxylase (putative GAD4) DNA From Arabidopsis genome sequencing project

Note: This is nucleic acid SEQ #7 and amino acid SEO #8

ACCESSION #AC006532

Part of chromosome #2

/product="putative glutamate decarboxylase" /protein_id="AAD20099.1" /db_xref="GI:4406789"

sis thaliana ecotype Columbia putativa alutama

5) Arabidopsis thaliana ecotype Columbia putative glutamate decarboxylase (putative GAD5) DNA From Arabidopsis genome sequencing project

Note: This is nucleic acid SEQ #9 and amino acid SEQ #10

ACCESSION #AB026646

Part of chromosome #3

/evidence=not_experimental

/product="glutamate decarboxylase" /protein_id="BAB02870.1" /db_xref="GI:9294589"

6) Tobacco (*Nicotiana tabacum*) glutamate decarboxylase isozyme 1 (NtGAD1) cDNA

Note: This is nucleic acid SEQ #11 and amino acid SEQ #12

A) LOCUS AF020425

ACCESSION AF020425

VERSION AF020425.1 GI:3252855

REFERENCE

AUTHORS Yun, S.J. and Oh, S.H.

TITLE Cloning and characterization of a tobacco cDNA encoding calcium/calmodulin-dependent glutamate decarboxylase JOURNAL Mol. Cells 8 (2), 125-129 (1998)

B) LOCUS NTU54774 ACCESSION U54774

VERSION U54774.1 GI:1777920

REFERENCE

AUTHORS Dharmasiri, M.A.N., Lu, Y.T. and Harrington, H.M.

TITLE Cloning and sequencing of a tobacco cDNA encoding glutamate decarboxylase

JOURNAL Unpublished

7) Tobacco (*Nicotiana tabacum*) glutamate decarboxylase isozyme 2 (NtGAD2) cDNA

Note: This is nucleic acid SEQ #13 and amino acid SEQ #14

LOCUS AF020424

ACCESSION AF020424

VERSION AF020424.1 GI:3252853

REFERENCE 1 (bases 1 to 1771)

AUTHORS Yun, S.J. and Oh, S.H.

TITLE Cloning and characterization of a tobacco cDNA encoding calcium/calmodulin-dependent glutamate decarboxylase

JOURNAL Mol. Cells 8 (2), 125-129 (1998)

8) Petunia (Petunia hybrida) glutamate decarboxylase cDNA

Note: This is nucleic acid SEQ #15 and amino acid SEQ #16

2. LOCUS PETGADX

ACCESSION #L16797

VERSION # L16797.1 GI:294111

KEYWORDS glutamate decarboxylase.

REFERENCE

AUTHORS Baum, G., Chen, Y., Arazi, T., Takatsuji, H. and Fromm, H.

TITLE A plant glutamate decarboxylase containing a calmodulin binding domain: cloning, sequence, and functional analysis

JOURNAL J. Biol. Chem. 268, 19610-19617 (1993)

B) LOCUS PETGLUDECA

ACCESSION L16977

VERSION L16977.1 GI:309679

REFERENCE

AUTHORS Baum, G., Chen, Y., Arazi, T., Takatsuji, H. and Fromm, H.

TITLE A plant glutamate decarboxylase containing a calmodulin-binding domain: cloning sequence and functional analysis

JOURNAL J. Biol. Chem. (1993)

9) Tomato (Lycopersicon esculentum) glutamate decarboxylase-like protein LEGDL cDNA

Note: This is nucleic acid SEQ #17 and amino acid SEQ #18

ACCESSION X80840

VERSION X80840.1 GI:993002

REFERENCE

AUTHORS Gallego, P.P., Whotton, L., Picton, S., Grierson, D. and Gray, J.E.

TITLE A role for glutamate decarboxylase during tomato ripening: the characterization of a cDNA encoding a putative glutamate decarboxylase with a calmodulin-binding site

JOURNAL Plant Mol. Biol. 27 (6), 1143-1151 (1995)